

DATE: 01/28/2003

1600

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TIME: 17:02:22
                     PATENT APPLICATION: US/09/677,653A
                     Input Set : A:\A-58631-4.txt
                     Output Set: N:\CRF4\01282003\1677653A.raw
                     SEQUENCE LISTING
      3 (1) GENERAL INFORMATION:
             (i) APPLICANT: P. D. CHRISTIAN, K. H. J. GORDON and
      5
                            T. N. HANZLIK
      7
            (ii) TITLE OF INVENTION: INSECT VIRUSES AND THEIR USES IN
                                      PROTECTING PLANTS
      8
     10
           (iii) NUMBER OF SEQUENCES: 53
     12
            (iv) CORRESPONDENCE ADDRESS:
                  (A) ADDRESSEE: DORSEY & WHITNEY LLP
     13
                  (B) STREET: FOUR EMBARCADERO CENTER, SUITE 3400
     14
     15
                  (C) CITY: SAN FRANCISCO
     16
                  (D) STATE: CALIFORNIA
     17
                  (E) COUNTRY: UNITED STATES OF AMERICA
     18
                  (F) ZIP: 94111
            (v) COMPUTER READABLE FORM:
                                                                     ENTERED
                  (A) MEDIUM TYPE: Floppy disk
     21
     22
                  (B) COMPUTER: IBM PC compatible
                  (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     23
     24
                  (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
     26
            (vi) CURRENT APPLICATION DATA:
C--> 27
                  (A) APPLICATION NUMBER: US/09/677,653A
C--> 28
                  (B) FILING DATE: 03-Oct-2000
     29
                  (C) CLASSIFICATION:
     31
          (viii) ATTORNEY/AGENT INFORMATION:
                  (A) NAME: RICHARD F. TRECARTIN
     32
     33
                  (B) REGISTRATION NUMBER: 31,801
     34
                  (C) REFERENCE/DOCKET NUMBER: A-58631-4/RFT/NBC
     36
            (ix) TELECOMMUNICATION INFORMATION:
     37
                  (A) TELEPHONE: (415) 781-1989
     40 (2) INFORMATION FOR SEQ ID NO: 1:
             (i) SEQUENCE CHARACTERISTICS:
     41
                  (A) LENGTH: 13 base pairs
     42
     43
                  (B) TYPE: nucleic acid
     44
                  (C) STRANDEDNESS: single
     45
                  (D) TOPOLOGY: linear
     47
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
     49 GGATCCACAG NNN
                                                                                 13
     51 (2) INFORMATION FOR SEQ ID NO: 2:
     52
             (i) SEQUENCE CHARACTERISTICS:
     53
                  (A) LENGTH: 28 base pairs
     54
                  (B) TYPE: nucleic acid
     55
                  (C) STRANDEDNESS: single
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RAW SEQUENCE LISTING

(D) TOPOLOGY: linear

56

## RAW SEQUENCE LISTING

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Input Set : A:\A-58631-4.txt

Output Set: N:\CRF4\01282003\I677653A.raw

```
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
58
60 ATGGGCGATG CCGGCGTCGC GTTCACAG
                                                                           28
62 (2) INFORMATION FOR SEQ ID NO: 3:
       (i) SEQUENCE CHARACTERISTICS:
64
             (A) LENGTH: 27 base pairs
             (B) TYPE: nucleic acid
66
             (C) STRANDEDNESS: single
             (D) TOPOLOGY: linear
67
       (xi) SEOUENCE DESCRIPTION: SEO ID NO: 3:
69
71 ATGGAGGATG CTGGAGTGGC GTCACAG
                                                                           27
73 (2) INFORMATION FOR SEQ ID NO: 4:
       (i) SEQUENCE CHARACTERISTICS:
75
             (A) LENGTH: 27 base pairs
76
             (B) TYPE: nucleic acid
77
             (C) STRANDEDNESS: single
78
             (D) TOPOLOGY: linear
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
82 ATGAGCGAGG CCGGCGTCGC GTCACAG
                                                                           27
84 (2) INFORMATION FOR SEQ ID NO: 5:
       (i) SEQUENCE CHARACTERISTICS:
86
             (A) LENGTH: 30 base pairs
             (B) TYPE: nucleic acid
87
             (C) STRANDEDNESS: single
88
89
             (D) TOPOLOGY: linear
91
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
                                                                           30
93 CCATCGATGC CGGACTGGTA TCCCAGGGGG
96 (2) INFORMATION FOR SEQ ID NO: 6:
        (i) SEQUENCE CHARACTERISTICS:
98
            (A) LENGTH: 31 base pairs
99
             (B) TYPE: nucleic acid
100
             (C) STRANDEDNESS: single
101
              (D) TOPOLOGY: linear
103
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
105 CCATCGATGC CGGACTGGTA TCCCGAGGGA C
                                                                            31
107 (2) INFORMATION FOR SEQ ID NO: 7:
108
        (i) SEQUENCE CHARACTERISTICS:
109
             (A) LENGTH: 39 base pairs
              (B) TYPE: nucleic acid
110
111
              (C) STRANDEDNESS: single
              (D) TOPOLOGY: linear
112
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
116 CCATCGATGA TCCAGCCTCC TCGCGGCGCC GGATGGGCA
                                                                            39
118 (2) INFORMATION FOR SEQ ID NO: 8:
119
        (i) SEQUENCE CHARACTERISTICS:
120
             (A) LENGTH: 39 base pairs
121
              (B) TYPE: nucleic acid
122
              (C) STRANDEDNESS: single
123
              (D) TOPOLOGY: linear
125 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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RAW SEQUENCE LISTING DATE: 01/28/2003 PATENT APPLICATION: US/09/677,653A TIME: 17:02:22

Input Set : A:\A-58631-4.txt

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127 GCTCTAGATC CATTCG	CCAT CCGAAGATGC CCATCCGGC	39
129 (2) INFORMATION F		33
	CHARACTERISTICS:	
· · · <del></del>	GTH: 39 base pairs	
	E: nucleic acid	
	ANDEDNESS: single	
	OLOGY: linear	
	DESCRIPTION: SEQ ID NO: 9:	
	AGAA GGTAACCAGA GAAACACAC	39
140 (2) INFORMATION F		33
	CHARACTERISTICS:	
	GTH: 39 base pairs	
• •	E: nucleic acid	
• •	ANDEDNESS: single	
• •	OLOGY: linear	
, , ,	DESCRIPTION: SEQ ID NO: 10:	
	TATA CCACAACGTG TGTTTCTCT	39
152 (2) INFORMATION F		39
	CHARACTERISTICS:	
· · · · · · · · · · · · · · · · · · ·		
	GTH: 45 base pairs E: nucleic acid	
	ANDEDNESS: single	
• •	OLOGY: linear	
	DESCRIPTION: SEQ ID NO: 11: GGTG ACACTATAGT TCTGCCTCCC CGGAC	45
163 (2) INFORMATION F		45
	CHARACTERISTICS:	
	GTH: 27 base pairs	•
, ,	E: nucleic acid	
, ,	ANDEDNESS: single	
	OLOGY: linear	
	DESCRIPTION: SEQ ID NO: 12:	
170 (XI) SEQUENCE		27
174 (2) INFORMATION F		21
	CHARACTERISTICS:	
` ' ~ · ~	GTH: 28 base pairs	
• •	E: nucleic acid	
· · ·	ANDEDNESS: single	
	OLOGY: linear	
	DESCRIPTION: SEQ ID NO: 13:	
183 CCGGAAGCTT GTTTTT		28
185 (2) INFORMATION F		20
	CHARACTERISTICS:	
<del>-</del>	GTH: 46 base pairs	
· •	E: nucleic acid	
	ANDEDNESS: single	
, ,	OLOGY: linear	
• •	DESCRIPTION: SEQ ID NO: 14:	
	TCCC GAGGGACGCT CAGCAGGTGG CATAGG	46
TO GOODATCO AIGGIA	ICCC GAGGGACGCI CAGCAGGIGG CAIAGG	40

RAW SEQUENCE LISTING DATE: 01/28/2003 PATENT APPLICATION: US/09/677,653A TIME: 17:02:22

Input Set : A:\A-58631-4.txt

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196 (2) INFORMATION FOR SEQ ID NO: 15:
    (i) SEQUENCE CHARACTERISTICS:
197
198
             (A) LENGTH: 52 base pairs
199
             (B) TYPE: nucleic acid
200
             (C) STRANDEDNESS: single
201
              (D) TOPOLOGY: linear
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
203
205 AAATAATTTT GTTACTTTAG AAGGAGATAT ACATATGAGC GAGCGAGCAC AC
                                                                          52
207 (2) INFORMATION FOR SEQ ID NO: 16:
    (i) SEQUENCE CHARACTERISTICS:
208
209
             (A) LENGTH: 55 base pairs
210
             (B) TYPE: nucleic acid
211
              (C) STRANDEDNESS: single
212
             (D) TOPOLOGY: linear
214
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
216 AAATAATTTT GTTTAACCTT AAGAAGGAGA TCTACATATG CTGGAGTGGC GTCAC
                                                                          55
218 (2) INFORMATION FOR SEQ ID NO: 17:
219 (i) SEQUENCE CHARACTERISTICS:
220
            (A) LENGTH: 30 base pairs
221
             (B) TYPE: nucleic acid
222
             (C) STRANDEDNESS: single
223
             (D) TOPOLOGY: linear
225 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
                                                                          30
227 GGAGATCTAC ATATGGGAGA TGCTGGAGTG
229 (2) INFORMATION FOR SEQ ID NO: 18:
230 (i) SEQUENCE CHARACTERISTICS:
231
            (A) LENGTH: 17 base pairs
232
             (B) TYPE: nucleic acid
233
             (C) STRANDEDNESS: single
234
             (D) TOPOLOGY: linear
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
236
                                                                          17
238 GTAGCGAACG TCGAGAA
240 (2) INFORMATION FOR SEQ ID NO: 19:
241 (i) SEQUENCE CHARACTERISTICS:
242
            (A) LENGTH: 31 base pairs
243
             (B) TYPE: nucleic acid
244
             (C) STRANDEDNESS: single
             (D) TOPOLOGY: linear
245
247
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
249 GGGGGATCCT CAGTTGTCAG TGGCGGGGTA G
                                                                          31
251 (2) INFORMATION FOR SEQ ID NO: 20:
252
      (i) SEQUENCE CHARACTERISTICS:
253
             (A) LENGTH: 28 base pairs
254
             (B) TYPE: nucleic acid
255
             (C) STRANDEDNESS: single
256
             (D) TOPOLOGY: linear
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
260 GGGGATCCCT AATTGGCACG AGCGGCGC
                                                                          28
263 (2) INFORMATION FOR SEQ ID NO: 21:
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## PATENT APPLICATION: US/09/677,653A TIME: 17:02:22 Input Set : A:\A-58631-4.txt Output Set: N:\CRF4\01282003\I677653A.raw 264 (i) SEQUENCE CHARACTERISTICS: 265 (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid 266 (C) STRANDEDNESS: single 267 268 (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21: 272 AATTACATAT GGCGGCCGCC GTTTCTGCC 29 274 (2) INFORMATION FOR SEQ ID NO: 22: (i) SEQUENCE CHARACTERISTICS: 276 (A) LENGTH: 29 base pairs 277 (B) TYPE: nucleic acid 278 (C) STRANDEDNESS: single (D) TOPOLOGY: linear 279 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22: 281 283 AATTACATAT GTTCGCGGCC GCCGTTTCT 29 285 (2) INFORMATION FOR SEQ ID NO: 23: 286 (i) SEQUENCE CHARACTERISTICS: 287 (A) LENGTH: 19 amino acids 288 (B) TYPE: amino acid (C) STRANDEDNESS: single 289 290 (D) TOPOLOGY: linear (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23: 292 Phe Ala Ala Ala Val Ser Ala Phe Ala Ala Asn Met Leu Ser Ser Val 294 295 1 10 296 Leu Lys Ser 299 (2) INFORMATION FOR SEQ ID NO: 24: (i) SEQUENCE CHARACTERISTICS: 301 (A) LENGTH: 20 amino acids 302 (B) TYPE: amino acid (C) STRANDEDNESS: single 303 304 (D) TOPOLOGY: linear 306 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24: 308 Pro Thr Leu Val Asp Gln Gly Phe Trp Ile Gly Gly Gln Tyr Ala Leu 309 1 310 Thr Pro Thr Ser 311 20 313 (2) INFORMATION FOR SEQ ID NO: 25: 314 (i) SEQUENCE CHARACTERISTICS: 315 (A) LENGTH: 6 amino acids 316 (B) TYPE: amino acid 317 (C) STRANDEDNESS: single 318 (D) TOPOLOGY: linear 320 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25: 322 Phe Ala Ala Val Ser 323 1 324 (2) INFORMATION FOR SEQ ID NO: 26: 326 (i) SEQUENCE CHARACTERISTICS: 327 (A) LENGTH: 23 base pairs 328 (B) TYPE: nucleic acid

RAW SEQUENCE LISTING

VERIFICATION SUMMARY

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PATENT APPLICATION: US/09/677,653A

TIME: 17:02:23

Input Set : A:\A-58631-4.txt

Output Set: N:\CRF4\01282003\I677653A.raw

L:27 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:28 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]